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PDB ID	Experimental Technique	Title	Authors	First Page	PubMed ID	Journal Name	Volume No	Publication Year
<u>1ALU</u>	X-RAY DIFFRACTION	HUMAN INTERLEUKIN-6	W.Somers, M.Stahl, J.S.Seehra	989	<u>9118960</u>	EMBO J	16	1997
<u>1BBN</u>	NMR	THREE-DIMENSIONAL SOLUTION STRUCTURE OF HUMAN INTERLEUKIN-4 BY MULTI-DIMENSIONAL HETERONUCLEAR MAGNETIC RESONANCE SPECTROSCOPY	R.Powers, D.S.Garrett, C.J.March, E.A.Frieden, A.M.Gronenborn, G.M.Clore	1673	<u>1609277</u>	Science	256	1992
<u>1BCN</u>	NMR	THREE-DIMENSIONAL SOLUTION STRUCTURE OF HUMAN INTERLEUKIN-4 BY MULTI-DIMENSIONAL HETERONUCLEAR MAGNETIC RESONANCE SPECTROSCOPY	R.Powers, D.S.Garrett, C.J.March, E.A.Frieden, A.M.Gronenborn, G.M.Clore	1673	<u>1609277</u>	Science	256	1992
<u>1BJ8</u>	NMR, MINIMIZED AVERAGE STRUCTURE	THIRD N-TERMINAL DOMAIN OF GP130, NMR, MINIMIZED AVERAGE STRUCTURE	T.Kernebeck, S.Pflanz, G.Muller-Newen, R.M.Scheek, K.Dijkstra, P.C.Heinrich, A.Wollmer, S.Grzesiek, J.Grotzinger	5	<u>10210178</u>	Protein Sci.	8	1999
<u>1BQU</u>	X-RAY DIFFRACTION	CYTOKINE-BINDING REGION OF GP130	J.Bravo, D.Staunton, J.K.Heath, E.Y.Jones	1665	<u>9501088</u>	EMBO J	17	1998
<u>1CYL</u>	NMR	ASPECTS OF RECEPTOR BINDING AND SIGNALLING OF INTERLEUKIN-4 INVESTIGATED BY SITE-	T.Muller, T.Dieckmann, W.Seibald, H.Oschkinat	423	<u>8151703</u>	J Mol Biol	237	1994



<u>1ILR</u>	X-RAY DIFFRACTION	SOLUTION CRYSTAL STRUCTURE OF THE INTERLEUKIN-1 RECEPTOR ANTAGONIST	A.M.Gronenborn H.A.Schreuder, J.M.Rondeau, C.Tardif, A.Soffientini, E.Sarubbi, A.Akeson, T.L.Bowlin, S.Yanofsky, R.W.Barrett	838	<u>7867645</u>	Eur J Biochem	227	1995
<u>1INR</u>	X-RAY DIFFRACTION	CYTOKINE SYNTHESIS	M.R. Walter, T.L.Nagabhushan	12118	<u>7547951</u>	Biochemistry	34	1995
<u>1IRL</u>	NMR, MINIMIZED AVERAGE STRUCTURE	THE SOLUTION STRUCTURE OF THE F42A MUTANT OF HUMAN INTERLEUKIN 2	H.R.Mott, B.S.Baines, R.M.Hall, R.M.Cooke, P.C.Driscoll, M.P.Weir, I.D.Campbell	979	<u>7723044</u>	J Mol Biol	247	1995
<u>1ITB</u>	X-RAY DIFFRACTION	TYPE-1 INTERLEUKIN-1 RECEPTOR COMPLEXED WITH INTERLEUKIN-1 BETA	G.P.Vigers, L.J.Anderson, P.Caffes, B.J.Brandhuber	190	<u>9062193</u>	Nature	386	1997
<u>1ITL</u>	NMR	HUMAN INTERLEUKIN 4: THE SOLUTION STRUCTURE OF A FOUR-HELIX-BUNDLE PROTEIN	L.J.Smith, C.Redfield, J.Boyd, G.M.Lawrence, R.G.Edwards, R.A.Smith, C.M.Dobson	899	<u>1569578</u>	J Mol Biol	224	1992
<u>1ITM</u>	NMR	ANALYSIS OF THE SOLUTION STRUCTURE OF HUMAN INTERLEUKIN 4 DETERMINED BY HETERONUCLEAR THREE- DIMENSIONAL NUCLEAR MAGNETIC RESONANCE TECHNIQUES	C.Redfield, L.J.Smith, J.Boyd, G.M.Lawrence, R.G.Edwards, C.J.Gershater, R.A.Smith, C.M.Dobson	23	<u>8145254</u>	J Mol Biol	238	1994
<u>1JLI</u>	NMR, MINIMIZED AVERAGE STRUCTURE	HUMAN INTERLEUKIN 3 (IL- 3) MUTANT WITH TRUNCATION AT BOTH N- AND C-TERMINI AND 14 RESIDUE CHANGES, NMR, MINIMIZED AVERAGE STRUCTURE	Y.Feng, B.K.Klein, C.A.McWherter	524	<u>8676386</u>	J Mol Biol	259	1996
	X-RAY	CRYSTAL STRUCTURE OF	S.G.Hymowitz,	5332	<u>11574464</u>	Embo J.	20	2001

<u>1JPY</u>	DIFFRACTION	IL-17F	E.H.Filvaroff, J.P. Yin, J.Lee, L.Cai, P.Risser, M.Maruoka, W.Mao, J.Foster, R.F.Kelley, G.Pan, A.L.Gurney, A.M.De Vos, M.A.Starovasnik						
<u>1JXQ</u>	X-RAY DIFFRACTION	STRUCTURE OF CLEAVED, CARD DOMAIN DELETED CASPASE-9	M.Renatus, H.R.Stennicke, F.L.Scott, R.C.Liddington, G.S.Salvesen	14250	<u>11734640</u>	Proc.Nat.Acad.Sci.USA	98	2001	
<u>1JXS</u>	NMR, 20 STRUCTURES	SOLUTION STRUCTURE OF THE DNA-BINDING DOMAIN OF INTERLEUKIN ENHANCER BINDING FACTOR	P.P.Liu, Y.C.Chen, C.Li, Y.H.Hsieh, S.W.Chen, S.H.Chen, W.Y.Jeng, W.J.Chuang	543	n/a	Proteins: Struct.,Funct.,Genet.	49	2002	
<u>1L2H</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF INTERLEUKIN 1-BETA F42W/W120F MUTANT	M.G.Rudolph, M.S.Kelker, T.R.Schneider, T.O.Yeates, V.Oseroff, D.K.Heidary, P.A.Jennings, I.A.Wilson	290	n/a	Acta Crystallogr., Sect.D	59	2003	
<u>1LOS</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF HUMAN CYTOMEGALOVIRUS IL-10 BOUND TO SOLUBLE HUMAN IL-10R1	B.C.Jones, N.J.Logsdon, K.Josephson, J.Cook, P.A.Barry, M.R.Walter	9404	<u>12093920</u>	Proc.Nat.Acad.Sci.USA	99	2002	
<u>1MD6</u>	X-RAY DIFFRACTION	HIGH RESOLUTION CRYSTAL STRUCTURE OF MURINE IL-1F5 REVEALS UNIQUE LOOP CONFORMATION FOR SPECIFICITY	E.F.Dunn, N.J.Gay, A.F.Bristow, D.P.Gearing, L.A.O'Neill, X.Y.Pei	10938	n/a	Biochemistry	42	2003	
<u>1N1F</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF HUMAN INTERLEUKIN-19	C.Chang, E.Magracheva, S.Kozlov, S.Fong, G.Tobin, S.Kotenko, A.Wlodawer, A.Zdanov	3308	n/a	J.Biol.Chem.	278	2003	

<u>1NAP</u>	X-RAY DIFFRACTION	THE CRYSTAL STRUCTURE OF RECOMBINANT HUMAN NEUTROPHIL-ACTIVATING PEPTIDE-2 (M6L) AT 1.9- ANGSTROMS RESOLUTION	M.G.Malkowski, J.Y.Wu, J.B.Lazar, P.H.Johnson, B.F.Edwards	7077	<u>7706245</u>	J Biol Chem	270	1995
<u>1O77</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF THE C713S MUTANT OF THE TIR DOMAIN OF HUMAN TLR2	X.Tao, Y.Xu, Y.Zheng, A.Beg, L.Tong	216	n/a	Biochem.Biophys.Res.Comm.	299	2002
<u>1OBX</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF THE COMPLEX OF PDZ2 OF SYNTENIN WITH AN INTERLEUKIN 5 RECEPTOR ALPHA PEPTIDE.	B.S.Kang, D.R.Cooper, Y.Devedjiev, U.Derewenda, Z.S.Derewenda	845	n/a	Structure	11	2003
<u>1OBY</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF THE COMPLEX OF PDZ2 OF SYNTENIN WITH A SYNDECAN-4 PEPTIDE.	B.S.Kang, D.R.Cooper, Y.Devedjiev, U.Derewenda, Z.S.Derewenda	845	n/a	Structure	11	2003
<u>1OBZ</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF THE COMPLEX OF THE PDZ TANDEM OF SYNTENIN WITH AN INTERLEUKIN 5 RECEPTOR ALPHA PEPTIDE.	B.S.Kang, D.R.Cooper, Y.Devedjiev, U.Derewenda, Z.S.Derewenda	845	n/a	Structure	11	2003
<u>1P9M</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF THE HEXAMERIC HUMAN IL-6/IL-6 ALPHA RECEPTOR/GP130 COMPLEX	M.J.Boulanger, D.C.Chow, E.E.Brevnova, K.C.Garcia	2101	n/a	Science	300	2003
<u>1PAU</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF THE COMPLEX OF APOPAIN WITH THE TETRAPEPTIDE ALDEHYDE INHIBITOR AC- DEVVD-CHO	J.Rotonda, D.W.Nicholson, K.M.Fazil, M.Gallant, Y.Gareau, M.Labelle, E.P.Peterson, D.M.Rasper, R.Ruel, J.P.Vaillancourt, N.A.Thornberry, J.W.Becker	619	<u>8673606</u>	Nat Struct Biol	3	1996
<u>1PVH</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF LEUKEMIA INHIBITORY FACTOR IN COMPLEX WITH GP130	M.J.Boulanger, A.J.Bankovich, T.Kortemme, D.Baker, K.C.Garcia	577	n/a	Mol. Cell	12	2003

<u>1PW6</u>	X-RAY DIFFRACTION	LOW MICROMOLAR SMALL MOLECULE INHIBITOR OF IL-2	C.D.Thanos, M.Randal, J.A.Wells	15280	n/a	J.Am.Chem.Soc.	125	2003
<u>1PY2</u>	X-RAY DIFFRACTION	STRUCTURE OF A 60 NM SMALL MOLECULE BOUND TO A HOT SPOT ON IL-2	C.D.Thanos, M.Randal, J.A.Wells	15280	n/a	J.Am.Chem.Soc.	125	2003
<u>1QDU</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF THE COMPLEX OF CASPASE-8 WITH THE TRIPEPTIDE KETONE INHIBITOR ZEVD-DCBMK	H.Blanchard, L.Kodandapani, P.R.E.Mittl, S.Di Marco, J.K.Krebs, J.C.Wu, K.J.Tomaselli, M.G.Grufter	1125	<u>10508784</u>	Structure (London)	7	1999
<u>1RCB</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF HUMAN RECOMBINANT INTERLEUKIN-4 AT 2.25 ANGSTROMS RESOLUTION	A.Wlodaver, A.Pavlovsky, A.Gustchina	59	<u>1511746</u>	FEBS Lett	309	1992
<u>1ROD</u>	NMR, 8 STRUCTURES	CHIMERIC PROTEIN OF INTERLEUKIN 8 AND HUMAN MELANOMA GROWTH STIMULATING ACTIVITY PROTEIN, NMR	H.Sticht, M.Auer, B.Schmitt, J.Besemer, M.Horcher, T.Kirsch, I.J.Lindley, P.Rosch	26	<u>8631339</u>	Eur J Biochem	235	1996
<u>1TVX</u>	X-RAY DIFFRACTION	NEUTROPHIL ACTIVATING PEPTIDE-2 VARIANT FORM M6L WITH FIVE ADDITIONAL AMINO TERMINAL RESIDUES (DSDLY)	M.G.Malkowski, J.B.Lazar, P.H.Johnson, B.F.Edwards	367	<u>9047370</u>	J Mol Biol	266	1997
<u>2CYK</u>	NMR	ASPECTS OF RECEPTOR BINDING AND SIGNALLING OF INTERLEUKIN-4 INVESTIGATED BY SITE- DIRECTED MUTAGENESIS AND NMR SPECTROSCOPY	T.Muller, T.Dieckmann, W.Seibald, H.Oschkinat	423	<u>8151703</u>	J Mol Biol	237	1994
<u>2FGF</u>	X-RAY DIFFRACTION	THREE-DIMENSIONAL STRUCTURE OF HUMAN BASIC FIBROBLAST GROWTH FACTOR, A STRUCTURAL HOMOLOG OF INTERLEUKIN 1BETA	J.D.Zhang, L.S.Cousens, P.J.Barr, S.R.Sprang	3446	<u>1849658</u>	Proc Natl Acad Sci U S A	88	1991
	X-RAY	CRYSTALLOGRAPHIC	J.P.Priestle, H.P.Schar,	9667	<u>2602367</u>	Proc Natl Acad Sci U S A	86	1989

<u>211B</u>	DIFFRACTION	REFINEMENT OF INTERLEUKIN-1BETA AT 2.0 ANGSTROMS RESOLUTION	M.G.Grutter					
<u>21L8</u>	NMR	THREE-DIMENSIONAL STRUCTURE OF INTERLEUKIN 8 IN SOLUTION	G.M.Clore, E.Appella, M.Yamada, K.Matsushima, A.M.Gronenborn	1689	<u>2184886</u>	Biochemistry	29	1990
<u>21LA</u>	X-RAY DIFFRACTION	STRUCTURE OF INTERLEUKIN 1ALPHA AT 2.7-ANGSTROMS RESOLUTION	B.J.Graves, M.H.Hatada, W.A.Hendrickson, J.K.Miller, V.S.Madison, Y.Satow	2679	<u>2346741</u>	Biochemistry	29	1990
<u>31L8</u>	X-RAY DIFFRACTION	CRYSTAL STRUCTURE OF INTERLEUKIN 8: SYMBIOSIS OF /NMR AND CRYSTALLOGRAPHY	E.T.Baldwin, I.T.Weber, R.St Charles, J.C.Xuan, E.Appella, M.Yamada, K.Matsushima, B.F.Edwards, G.M.Clore, A.M.Gronenborn, et al.	502	<u>1988949</u>	Proc Natl Acad Sci U S A	88	1991
<u>31NK</u>	X-RAY DIFFRACTION	UNRAVELING THE STRUCTURE OF INTERLEUKIN-2: REPLY	D.B.Mckay	412	n/a	Science	257	1992
<u>411B</u>	X-RAY DIFFRACTION	FUNCTIONAL IMPLICATIONS OF INTERLEUKIN-1BETA BASED ON THE THREE- DIMENSIONAL STRUCTURE	B.Veerapandian, G.L.Gilliland, R.Raag, A.L.Svensson, Y.Masui, Y.Hirai, T.L.Poulos	10	<u>1553379</u>	Proteins	12	1992
<u>511B</u>	X-RAY DIFFRACTION	A COMPARISON OF THE HIGH RESOLUTION STRUCTURES OF HUMAN AND MURINE INTERLEUKIN-1B	D.H.Ohlendorf, A.C.Trehanne, P.C.Weber, J.J.Wendoloski, F.R.Salemme, M.Lischwe, R.C.Newton	n/a	n/a	To be Published	n/a	n/a
<u>611B</u>	NMR	HIGH-RESOLUTION THREE- DIMENSIONAL STRUCTURE OF INTERLEUKIN 1BETA IN SOLUTION BY THREE-AND FOUR-DIMENSIONAL	G.M.Clore, P.T.Wingfield, A.M.Gronenborn	2315	<u>2001363</u>	Biochemistry	30	1991







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